	umber: CRF Errors Corrected by the STIC Systems Branch CRF Processing Date: Edited by:
	Changed a file from non-ASCII to ASCII Verified by: Verified by:
	Changed the margins in cases where the sequence text was "wrapped" down to the next line.
	Edited a format error in the Current Application Data section, specifical NTERED
	Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other
	Added the mandatory heading and subheadings for "Current Application Data".
/	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integ
	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
	Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included:
•	Deleted extra, invalid, headings used by an applicant, specifically:
	Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end ☐ page numbers throughout text; ☐ other invalid text, such as
	Inserted mandatory headings, specifically:
	Corrected an obvious error in the response, specifically:
	Edited identifiers where upper case is used but lower case is required, or vice versa.
/	Corrected an error in the Number of Sequences field, specifically:
	A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
,	Deleted <i>ending</i> stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (due to a PatentIn bug). Sequences corrected:
	Other:

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,913

DATE: 03/08/96 TIME: 16:38:10

INPUT SET: S9193.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

1				SEQUENCE LISTING	ENTERED
2 3 4	(1) Ge	eneral	Information:		
5 6	(i)	APPL	ICANT:	Andrew C. Hiatt	
7				Floyd Rose	
8 9 10 11 12 13	(ii)	TITL	E OF INVENTION:	3' PROTECTED NUC FOR ENZYME CATAL TEMPLATE-INDEPEN OF PHOSPHODIESTE BONDS	YZED DENT CREATION
14 15	(iii)	NUMB	ER OF SEQUENCES:	8	
16 17	(iv)	CORR	ESPONDENCE ADDRESS	5:	
18 19		(3)	ADDDDCCDD.	Twon C Twon	
20 21		(A) (B)	ADDRESSEE: STREET:	Lyon & Lyon 633 West Fifth S [.] Suite 4700	treet
22		(C)	CITY:	Los Angeles	
23		(D)	STATE:	California	
24		(E)	COUNTRY:	U.S.A.	
25		(F)	ZIP:	90071	
26		GOVE	ummo peanante con		
27 28	(V)	COMP	UTER READABLE FORM	1:	
29 30		(A)	MEDIUM TYPE:	3.5" Diskette, 1 storage	.44 Mb
31		(B)	COMPUTER:	IBM Compatible	
32		(C)	OPERATING SYSTEM	1: IBM P.C. DOS 5.0	
33		(D)	SOFTWARE:	Word Perfect 5.1	
34					•
35	(Vi)	CURR	ENT APPLICATION DA	ATA:	
36 37		/ 3 \	ADDITOAMTON NUMBE	ID.	
38		(A) (B)	APPLICATION NUMBE FILING DATE:	.R:	
39		(C)	CLASSIFICATION:		
40		(0)			
41	(vii)	PRIO	R APPLICATION DATA	\:	
42	. ,		r applications tot	al,	
43			uding application		
44		desc	ribed below:		
45	/*****	3 mm ^	DMDU/Addim TMDASYA	MTON.	
46	(viii)	ATTO	RNEY/AGENT INFORMA	TION:	

99

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,913

DATE: 03/08/96 TIME: 16:38:12

INPUT SET: S9193.raw (A) NAME: Guise, Jeffrey W. 47 (B) REGISTRATION NUMBER: 48 34,613 (C) REFERENCE/DOCKET NUMBER: 49 207/145 50 (ix) TELECOMMUNICATION INFORMATION: 51 52 53 (A) TELEPHONE: (619) 552-8400 54 (B) TELEFAX: (619) 552-0159 55 (C) TELEX: 67-3510 56 57 (2) INFORMATION FOR SEQ ID NO: 58 59 (i) SEQUENCE CHARACTERISTICS: 60 61 (A) LENGTH: 12 (B) TYPE: nucleic acid 62 63 (C) STRANDEDNESS: single (D) TOPOLOGY: linear 64 65 66 (ix) FEATURE: 67 68 (D) OTHER INFORMATION: base number 12 is m7g 69 70 71 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1: 72 73 CCCCCCCCC CG 12 74 75 76 77 (2) INFORMATION FOR SEQ ID NO: 78 (i) SEQUENCE CHARACTERISTICS: 79 80 81 (A) LENGTH: 19 82 (B) TYPE: nucleic acid 83 (C) STRANDEDNESS: single (D) TOPOLOGY: 84 linear 85 86 (ii) SEQUENCE DESCRIPTION: SEQ ID NO: 2: 87 CCCCCCCC CCCCTGCA 19 88 89 90 91 (2) INFORMATION FOR SEQ ID NO: 92 93 94 (i) SEQUENCE CHARACTERISTICS: 95 96 (A) LENGTH: 20 97 (B) TYPE: nucleic acid 98 (C) STRANDEDNESS: single

linear

(D) TOPOLOGY:

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,913

DATE: 03/08/96 TIME: 16:38:14

INPUT SET: S9193.raw

		HVF
100 101	(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 3:	
102	(II) bigoinel bibonititon, big is not st	
103	CTGCAGGGG GGGGGGGG	20
104		
105		
106	(2) INFORMATION FOR SEQ ID NO: 4:	
107		
108	(i) SEQUENCE CHARACTERISTICS:	
109	() Trucomit	
110 111	(A) LENGTH: 9 (B) TYPE: nucleic acid	
111	(B) TYPE: nucleic acid (C) STRANDEDNESS: single	
113	(D) TOPOLOGY: linear	
114	(5) 101 011001.	
115	(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 4:	
116	(
117	ccccccc	9
118		
119		
120	(2) INFORMATION FOR SEQ ID NO: 5:	
121		
122	(i) SEQUENCE CHARACTERISTICS:	
123 124	(A) LENGTH: 9	
124	(A) LENGTH: 9 (B) TYPE: nucleic acid	
126	(C) STRANDEDNESS: single	
127	(D) TOPOLOGY: linear	
128	(2) 101020017 1111001	
129	(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 5:	
130		
131	GGGGGGGG	9
132		
133		
134	(2) INFORMATION FOR SEQ ID NO: 6:	
135	(i) GEOLENGE GUADAGERIGETAG.	
136 137	(i) SEQUENCE CHARACTERISTICS:	
138	(A) LENGTH: 5	•
139	(B) TYPE: nucleic acid	
140	(C) STRANDEDNESS: single	
141	(D) TOPOLOGY: linear	
142	,	
143	(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 6:	
144		
145	CTGCA	5
146		
147	(2) INFORMATION FOR SEQ ID NO: 7:	
148		
149	(i) SEQUENCE CHARACTERISTICS:	
150	/A) I ENGMU. O	
151 152	(A) LENGTH: 9 (B) TYPE: nucleic acid	
127	(p) like: uncletc gcld	

\mathbf{p}_{Δ}	GF:	4
$r_{\mathcal{M}}$	UĽ.	+

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,913

DATE: 03/08/96 TIME: 16:38:17

INPUT SET: S9193.raw

		INPUL SEL: SYLYS.raw
153	(C) STRANDEDNESS: single	
154	(D) TOPOLOGY: linear	
155		
156	(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 7:	
157		
158	CTGCATGCA	9
159		
160		
161	(2) INFORMATION FOR SEQ ID NO: 8:	
162		
163	(i) SEQUENCE CHARACTERISTICS:	
164		
165	(A) LENGTH: 10	
166	(B) TYPE: nucleic acid	
167	(C) STRANDEDNESS: single	
168	(D) TOPOLOGY: linear	
169		
170	(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 8:	
171		
172	CTGCATGCAG	10
173		
174		
175		
176		
177		

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,913

DATE: 03/08/96 TIME: 16:38:18

INPUT SET: S9193.raw

***** PREVIOUSLY ERRORED SEQUENCES - EDITED *****

161	(2) INFORMATION FOR SEQ ID NO: 8:	
162	•	
163	(i) SEQUENCE CHARACTERISTICS:	
164		
165	(A) LENGTH: 10	
166	(B) TYPE: nucleic acid	
167	(C) STRANDEDNESS: single	
168	(D) TOPOLOGY: linear	
169		
170	(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 8:	
171		
172	CTGCATGCAG	10
173		
174		
175		
176		
177		

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/486,913

DATE: 03/08/96 TIME: 16:38:19

INPUT SET: S9193.raw

Line

Error

Original Text

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,913

DATE: 03/08/96 TIME: 09:50:50

INPUT SET: S9193.raw

This Raw Listing contains the General Information Section and those Sequences containing ERRORS.

	1			S	SEQUENCE LISTING	
	2					
	3	(1) Ge	neral	Information:		
	4					
	5 6	(i)	APPL	ICANT:	Andrew C. Hiatt	Does Not Comply
	7				Floyd Rose	Corrected Diskette Needed
	8				rioya Rose	
	9	(ii)	TTT.	E OF INVENTION:	3' PROTECTED NUCI	FOTINES
	10	(++)	1111	E OF INVENTION:	FOR ENZYME CATALY	
	11				TEMPLATE-INDEPEN	
	12				OF PHOSPHODIESTER	
	13				BONDS	•
	14					
>	15	(iii)	NUMB	ER OF SEQUENCES:	8	
	16	` •		~		
	17	(iv)	CORR	ESPONDENCE ADDRESS:		
	18					•
	19		(A)	ADDRESSEE:	Lyon & Lyon	
	20		(B)	STREET:	633 West Fifth St	reet
	21				Suite 4700	
	22		(C)	CITY:	Los Angeles	
	23		(D)	STATE:	California	
	24		(E)	COUNTRY:	U.S.A.	
	25		(F)	ZIP:	90071	
	26					
	27	(V)	COMP	UTER READABLE FORM:		
	28					
	29		(A)	MEDIUM TYPE:	3.5" Diskette, 1.	.44 Mb
	30				storage	
	31		(B)	COMPUTER:	IBM Compatible	
	32		(C)	OPERATING SYSTEM:		
	33		(D)	SOFTWARE:	Word Perfect 5.1	
	34 25	/ d \	GUDD.	ENE ADDITONEDAMA		
	35 36	(Vi)	CURR	ENT APPLICATION DATA	. .	
	36 37		(A)	APPLICATION NUMBER:		
	38		(A) (·B)	FILING DATE:		
	39		(C)	CLASSIFICATION:		
	40		(0)	CHADSIF ICATION:		
	41	(Vii)	PRTO	R APPLICATION DATA:		•
	42	(+ ± ± /		r applications total		
	43			uding application	• ,	
	44			ribed below:		
	45					

RAW SEQUENCE LISTING PATENT APPLICATION US/08/486,913

DATE: 03/08/96 TIME: 09:50:53

INPUT SET: S9193.raw

```
(viii)
               ATTORNEY/AGENT INFORMATION:
46
47
                    NAME:
                                                 Guise, Jeffrey W.
               (A)
48
               (B)
                    REGISTRATION NUMBER:
                                                 34,613
                    REFERENCE/DOCKET NUMBER:
                                                 207/145
49
               (C)
50
51
         (ix)
               TELECOMMUNICATION INFORMATION:
52
53
                     TELEPHONE:
                                                 (619) 552-8400
               (A)
54
               (B)
                    TELEFAX:
                                                 (619) 552-0159
                    TELEX:
                                                 67-3510(2)
                                                              INFORMATION FOR SEQ ID NO:
55
               (C)
56
                    SEQUENCE CHARACTERISTICS:
57
               (i)
58
59
                     (A)
                           LENGTH:
                                            nucleic acid
                     (B)
                           TYPE:
60
                     (C)
                           STRANDEDNESS:
                                            single
61
                           TOPOLOGY:
                                            linear
62
                     (D)
63
64
              (ix)
                    Features:
65
                     (D)
                           Other Information: base number 12 is m7g
66
67
68
                    SEQUENCE DESCRIPTION: SEQ ID NO: 1:
69
              (xi)
70
71
    CCCCCCCCC CG
                                                                12
72
73
74
```

ERRORED SEQUENCES FOLLOW:

			///
	159	(2) INFORMATION FOR SEQ ID NO: 8:	
	160		X
	161	(i) SEQUENCE CHARACTERISTICS:	()
	162		
	163	(A) LENGTH: 10	
	164	(B) TYPE: nucleic acid /	
	165	(C) STRANDEDNESS: single	
	166	(D) TOPOLOGY: linear	
	167		
>	168	(ii) SEQUENCE DESCRIPTION: SEQ ID NO: (10)	
	169		
	170	CTGCATGCAG 10	
	171		
	172		
	173		
	174		
	175		

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SEQUENCE VERIFICATION REPORT PATENT APPLICATION *US/08/486,913*

DATE: 03/08/96 TIME: 09:50:57

INPUT SET: S9193.raw

Line	Error	Original Text
15 57 59 60 61 62 64 66	Number of Sequences (8) Doesn't Equal Actual Count (7) Unknown or Misplaced Identifier	 (iii) NUMBER OF SEQUENCES: 8 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ix) Features: (D) Other Information: base number 12 is m7g (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
168	Wrong Sequence Number	(ii) SEQUENCE DESCRIPTION: SEQ ID NO: 10: